

## SEQUENCE LISTING

<110>	Lamberty, Mireille Bulet, Phillipe Brookhart, Gary Hoffman, Jules	
<120>	GENE CODING FOR HELIOMICINE, AND USE THEREOF	
<130>	A33595-PCT-USA (075188.0110)	
	09/673,274 1999-04-12	
	PCT/FR99/00843 1999-04-12	
	FR 98 04933 1998-04-15	
<160>	53	
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<210> <211> <212> <213>	147	
<220> <223>	Synthetic oligonucleotide	
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<210><211><211><212><213>	169	
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<220> <223>	Synthetic oligonucleotide	

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ttttccttgt gatctctcac tcttgccgtg ccgataagct tatcggttcc tgcgtgtggg 120
gtgctgtgaa ctacacttcc gattgcaacg gtgagtgcaa gaggaggggt tacaagggtg 180
gtcactgcgg ttccttcgct aacgtgaact gctggtgcga gacttgagag ctcggcgagg 240
cgaacgtgtc gacggatccg g
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gegtegaege gatgggttte gtgettttet eteagettee atettteett ettgtgteta 60
ctcttcttct tttccttgtg atctctcact cttgccgtgc tggagacgcg aattcacaca 120
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ctcttcttct tttcc
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aaagatggaa gc
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gataagetta teggtteetg egtgtggggt getgtgaaet acaetteega ttgcaaeggt 60
gagtgcaaga ggaggggtta
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<211> 109
<212> DNA
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ceggateegt egacaegtte geetegeega geteteaagt etegeaceag eagtteaegt 60
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agggcccct agggtttaaa cggccagtca ggccgaattc gagctcggta cccggggatc 60
ctctagagtc gacctgcagg catgc
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<211> 66
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<223> Synthetic oligonucleotide
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ccctgaacca ggctcgaggg cgcgccttaa ttaaaagctt gcatgcctgc aggtcgactc 60
tagagg
<210> 11
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ccggccagtc aggccacact taattaagtt taaacgcggc cccggcgcgc ctaggtgtgt 60
gctcgagggc ccaacctcag tacctggttc agg
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<223> Synthetic oligonucleotide
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gcgtttaaac ttaattaagt gtggcctgac tgg
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<400> 14
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ccgaattcgg cgccgtgcac gatgcagaag agcacgagga ggaagagggc
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<223> Synthetic oligonucleotide
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totagaatgg cotgoaccaa caacgocatg agggocotot tootcotoot gotottotgc 60
atcgtgcacg gcgccgaatt c
<210> 16
<211> 24
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gataagctta tcggttcctg cgtg
                                                                    24
<210> 17
<211> 32
<212> DNA
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ggctcgagtc aagtctcgca ccagcagttc ac
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<210> 18
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<212> DNA
<213> Artificial Sequence
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totagaatgg cotgoaccaa caacgocatg agggocotot tootootoot gotottotgc 60
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gattgcaacg gtgagtgcaa gaggagggt tacaagggtg gtcactgcgg ttccttcgct 180
aacgtgaact gctggtgcga gacttgactc gag
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<221> promoter
<222> (7)...(532)
<221> misc structure
<222> (533)...(568)
<221> terminator
<222> (569)...(832)
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actatggaag tattatgtga gctcagcaag aagcagatca atatgcggca catatgcaac 120
ctatgttcaa aaatgaagaa tgtacagata caagatccta tactgccaga atacgaagaa 180
gaatacgtag aaattgaaaa agaagaacca ggcgaagaaa agaatcttga agacgtaagc 240
actgacgaca acaatgaaaa gaagaagata aggtcggtga ttgtgaaaga gacatagagg 300
acacatgtaa ggtggaaaat gtaagggcgg aaagtaacct tatcacaaag gaatcttatc 360
ccccactact tatcctttta tatttttccg tgtcattttt gcccttgagt tttcctatat 420
aaggaaccaa gttcggcatt tgtgaaaaca agaaaaaatt tggtgtaagc tattttcttt 480
gaagtactga ggatacaact tcagagaaat ttgtaagttt gtagatctcg attctagaag 540
gcctgaattc gagctcggta ccggatccaa ttcccgatcg ttcaaacatt tggcaataaa 600
gtttcttaag attgaatcct gttgccggtc ttgcgatgat tatcatataa tttctgttga 660
attacgttaa gcatgtaata attaacatgt aatgcatgac gttatttatg agatgggttt 720
ttatgattag agtcccgcaa ttatacattt aatacgcgat agaaaacaaa atatagcgcg 780
caaactagga taaattatcg cgcgcggtgt catctatgtt actagatcgg ggatcgat
<210> 20
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<221> promoter
<222> (7)...(532)
<221> CDS
<222> (539)...(736)
<221> terminator
<222> (767)...(1030)
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aagcttccag aaggtaatta tccaagatgt agcatcaaga atccaatgtt tacgggaaaa 60
actatggaag tattatgtga gctcagcaag aagcagatca atatgcggca catatgcaac 120
ctatgttcaa aaatgaagaa tgtacagata caagatccta tactgccaga atacgaagaa 180
gaatacgtag aaattgaaaa agaagaacca ggcgaagaaa agaatcttga agacgtaagc 240
actgacgaca acaatgaaaa gaagaagata aggtcggtga ttgtgaaaga gacatagagg 300
acacatgtaa ggtggaaaat gtaagggcgg aaagtaacct tatcacaaag gaatcttatc 360
ccccactact tatcctttta tatttttccg tgtcattttt gcccttgagt tttcctatat 420
aaggaaccaa gttcggcatt tgtgaaaaca agaaaaaatt tggtgtaagc tattttcttt 480
gaagtactga ggatacaact tcagagaaat ttgtaagttt gtagatctcg attctaga
atg gcc tgc acc aac acc gcc atg agg gcc ctc ttc ctc ctc gtg ctc
                                                                   586
Met Ala Cys Thr Asn Asn Ala Met Arg Ala Leu Phe Leu Leu Val Leu
1
                                     10
                                                          15
                                                                   634
tte tge ate gtg cae gge gat aag ett ate ggt tee tge gtg tgg ggt
Phe Cys Ile Val His Gly Asp Lys Leu Ile Gly Ser Cys Val Trp Gly
             20
                                 25
gct gtg aac tac act tcc gat tgc aac ggt gag tgc aag agg agg ggt
                                                                   682
Ala Val Asn Tyr Thr Ser Asp Cys Asn Gly Glu Cys Lys Arg Arg Gly
         35
                             40
tac aag ggt ggt cac tgc ggt tcc ttc gct aac gtg aac tgc tgg tgc
                                                                   730
Tyr Lys Gly Gly His Cys Gly Ser Phe Ala Asn Val Asn Cys Trp Cys
     50
                         55
                                                                   786
gag act tgactcgagg gggggcccgg taccggatcc aattcccgat cgttcaaaca
Glu Thr
 65
tttggcaata aagtttctta agattgaatc ctgttgccgg tcttgcgatg attatcatat 846
aatttctgtt gaattacgtt aagcatgtaa taattaacat gtaatgcatg acgttattta 906
tgagatgggt ttttatgatt agagtcccgc aattatacat ttaatacgcg atagaaaaca 966
aaatatagcg cgcaaactag gataaattat cgcgcgcggt gtcatctatg ttactagatc 1026
ggggatcgat
                                                                   1036
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<210> 22
<211> 56
<212> DNA
<213> Artificial Sequence
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agtgtagttg acggcgcccc aaacacagct gccaatcaac ttgtctcttt tatcca
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<210> 23
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<211> 52
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<223> Synthetic oligonucleotide
<400> 23
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actacactag tgactgcaac ggcgagtgca agcgccgcgg ttacaagggt gg
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<211> 52
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide
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cacaatggcc accettgtaa eegeggeget tgcactegee gttgcagtca et
                                                                    52
<210> 25
<211> 56
<212> DNA
<213> Artificial Sequence
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<223> Synthetic oligonucleotide
ccattgtgga tccttcgcta acgttaactg ttggtgtgaa acctgatagg tcgaca
                                                                    56
<210> 26
<211> 52
<212> DNA
<213> Artificial Sequence
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<223> Synthetic oligonucleotide
<400> 26
gatctgtcga cctatcaggt ttcacaccaa cagttaacgt tagcgaagga tc
                                                                    52
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<211> 42
<212> DNA
<213> Artificial Sequence
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<223> Synthetic oligonucleotide
gatccttcgc taacgttaac tgttggtgta gaacctgata gg
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<210> 28
<211> 42
<212> DNA
<213> Artificial Sequence
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<400> 28 tcgaccta	3 atc aggttctaca ccaacagtta	acgttagcga	ag	42
<210> 29 <211> 32 <212> DN <213> Ar	2			
<220> <223> Sy	ynthetic oligonucleotide			
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agcttggata aaagagctgc tgctgctggt agctgtgttt	40
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<400> 34 ggggcgccgt caactaca	18
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<210> 36 <211> 36 <212> DNA <213> Artificial Sequence	
<220> <223> Synthetic oligonucleotide	
<400> 36 aaacacagct accagcagca gcagctcttt tatcca	36
<210> 37 <211> 32 <212> DNA <213> Artificial Sequence	
<220> <223> Synthetic oligonucleotide	
<400> 37 ctagtgactg cgctgctgag tgcttgttgc gc	32
<210> 38 <211> 26 <212> DNA <213> Artificial Sequence	
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<400> 38 gcaacaagca ctcagcagcg cagtca	26
<210> 39 <211> 51	

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<212> PRT
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<221> VARIANT
<222> (1)...(10)
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      where Xaa = any amino acid
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<222> (12)...(21)
<223> region of variable length from 1 to 10 amino acids
      where Xaa = any amino acid
<221> VARIANT
<222> (23)...(25)
<223> Xaa = any amino acid
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<222> (27)...(35)
<223> region of variable length from 1 to 9 amino acids
      where Xaa = any amino acid
<221> VARIANT
<222> (37)...(43)
<223> region of variable length from 1 to 7 amino acids
      where Xaa = any amino acid
<221> VARIANT
<222> (45)...(45)
<223> Xaa = any amino acid
<221> VARIANT
<222> (47)...(51)
<223> region of variable length from 1 to 5 amino acids
      where Xaa = any amino acid
<400> 39
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa
Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa
Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Cys Xaa Xaa
Xaa Xaa Xaa
    50
<210> 40
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<222> (2)...(2)
<223> Xaa = any basic amino acid
<221> VARIANT
<222> (3)...(7)
<223> region of variable length from 0 to 5 amino acids
      where Xaa = any amino acid
<221> VARIANT
<222> (1)...(9)
<223> Xaa = Any Amino Acid
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Lys Xaa Xaa Xaa Xaa Xaa Gly His
                 5
<210> 41
<211> 9
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<220>
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Lys Arg Arg Gly Tyr Lys Gly Gly His
<210> 42
<211> 11
<212> PRT
<213> Artificial Sequence
<223> Synthetic peptide
<221> VARIANT
<222> (1)...(9)
<223> region of variable length from 0 to 9 amino acids
     where Xaa = any amino acid
<221> VARIANT
<222> (11)...(11)
<223> Xaa = any amino acid
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<222> (1)...(11)
<223> Xaa = Any Amino Acid
<400> 42
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Gly Xaa
<210> 43
<211> 10
<212> PRT
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<213> Artificial Sequence
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<223> Synthetic peptide
<221> VARIANT
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      where Xaa = any amino acid
<221> VARIANT
<222> (1)...(10)
<223> Xaa = Any Amino Acid
<400> 43
Val Xaa Xaa Xaa Xaa Xaa Xaa Asp
1
                 5
<210> 44
<211> 7
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic peptide
<221> VARIANT
<222> (2)...(6)
<223> region of variable length from 0 to 5 amino acids
      where Xaa = any amino acid
<221> VARIANT
<222> (1)...(7)
<223> Xaa = Any Amino Acid
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Gly Xaa Xaa Xaa Xaa Asn
1
                 5
<210> 45
<211> 5
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<223> Synthetic peptide
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      where Xaa = any amino acid
<221> VARIANT
<222> (1)...(5)
<223> Xaa = Any Amino Acid
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Val Trp Gly Ala Val Asn Tyr Thr Ser Asp
<210> 48
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<213> Artificial Sequence
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<400> 48
Gly Ser Phe Ala Asn Val Asn
1
<210> 49
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<223> Synthetic peptide
<400> 49
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1
<210> 50
<211> 3
<212> PRT
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Glu Xaa Xaa Xaa Xaa

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<213> Artificial Sequence
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<223> Synthetic peptide
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Ala Ala Ala Gly Ser
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